SEQ ID NO:42	SEQ ID NO:43	SEQ ID NO:58	SEQ ID NO:66	SEQ ID NO:67	SEO ID NO:68
GNLTSQNVTTVTQPLKKTKS	GNLTSQNVTTVSPPLKKTKS	GSLTVNPKAPLQVNTD	GKLTVNTEPPLHLTNN	GKLTVNADPPLQLTNN	GNLTVNTEPPLOLTNN
Ad2	Ad5	Ad37	Ad8	Ad9	Ad15

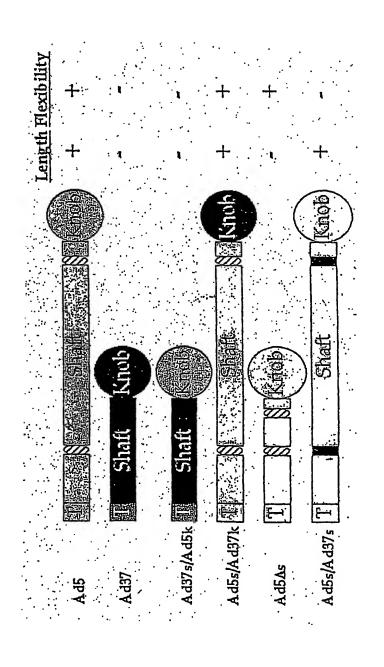
Alignment of the Corresponding Third Repeats of Ad Fiber Shaft Proteins

Figure 1A

	N	
	1S KLGXGLXFD	Consensus
SEQ ID NO:61	VRVGEGGGLSFNEA	Ad15
SEQ ID NO:60	VRVGEGGGLSFNND	Ad9
SEQ ID NO:59	VRVGEGGGLSFNDN	Ad8
SEQ ID NO:48	IGINVRAREGLTFDND	Ad37
SEQ ID NO:47	KAMVPKLGTGLSFDST	Ad5
SEQ ID NO:46	GAMITKLGAGLSFDNS	Ad2

Corresponding Last Full Fiber Shaft Proteins Alignment of the Ad of Repeats

Figure 1B



TGURE